# Figure 1-A

>2767646	GCCAGGAATA	ACTAGAGAGG	AACAATGGGG	TTATTCAGAG	GTTTTGTTTT
>774134		CTAGAGAGG	AACAATGGGG	TTATTCAGAG	GTTTTGTTTT
>774134IH		CTAGAGAGG	AACAATGGGG	TTATTCAGAG	GTTTTGTTTT
Consensus	GCCAGGAATA	ACTAGAGAGG	AACAATGGGG	TTATTCAGAG	GTTTTGTTTT
>2767646	CCTCTTAGTT	CTGTGCCTGC	TGCACCAGTC	AAATACTTCC	TTCATTAAGC
>774134	CCTCTTAGTT	CTGTGCCTGC	TGCACCAGTC	AAATACTTCC	TTCATTAAGC
>774134IH	CCTCTTAGTT	CTGTGCCTGC	TGCACCAGTC	AAATACTTCC	TTCATTAAGC
Consensus	CCTCTTAGTT	CTGTGCCTGC	TGCACCAGTC	AAATACTTCC	TTCATTAAGC
>2767646	TGAATAATAA	TGGCTTTGAA	GATATTGTCA	TTGTTATAGA	TCCTAGTGTG
>774134	TGAATAATAA	TGGCTTTGAA	GATATTGTCA	TTGTTATAGA	TCCTAGTGTG
>774134IH	TGAATAATAA	TGGCTTTGAA	GATATTGTCA	TTGTTATAGA	TCCTAGTGTG
Consensus	TGAATAATAA	TGGCTTTGAA	GATATTGTCA	TTGTTATAGA	TCCTAGTGTG
>2767646	CCAGAAGATG	ТААТААААА	TGAACAAATA	GAGGATATGG	TGACTACAGC
>774134	CCAGAAGATG	AAAAAATAAT	TGAACAAATA	GAGGATATGG	TGACTACAGC
>774134IH	CCAGAAGATG	${\tt AAAAAATAAT}$	TGAACAAATA	GAGGATATGG	TGACTACAGC
Consensus	CCAGAAGATG	TAAAAAAT	TGAACAAATA	GAGGATATGG	TGACTACAGC
>2767646	TTCTACGTAC	CTGTTTGAAG	CCACAGAAAA	AAGATTTTTT	т
>774134		CTGTTTGAAG			
>774134IH			CCACAGAAAA		
Consensus	TTCTACGTAC	CTGTTTGAAG	CCACAGAAAA	AAGATTTTTT	TTCAAAAATG
>774134IH			AATTGGAAGG		
Consensus	TATCTATATT	AATTCCTGAG	AATTGGAAGG	AAAATCCTCA	GTACAAAAGG
>774134IH			ACATGCTGAT		
Consensus	CCAAAACATG	AAAACCATAA	ACATGCTGAT	GTTATAGTTG	CACCACCTAC
>774134IH	ACTCCCAGGT	AGAGATGAAC	CATACACCAA	GCAGTTCACA	GAATGTGGAG
Consensus	ACTCCCAGGT	AGAGATGAAC	CATACACCAA	GCAGTTCACA	GAATGTGGAG
>774134IH			TTCACCCCTG		
Consensus	AGAAAGGCGA	ATACATTCAC	TTCACCCCTG	ACCTTCTACT	TGAAAAAAA
>774134IH	CAAAATGAAT	ATGGACCACC	AGGCAAACTG	TTTGTCCATG	AGTGGGCTCA
Consensus	CAAAATGAAT	ATGGACCACC	AGGCAAACTG	TTTGTCCATG	AGTGGGCTCA
>774134IH			ATGAGTACAA		
Consensus	CCTCCGGTGG	GGAGTGTTTG	ATGAGTACAA	TGAAGATCAG	CCTTTCTACC
>774134IH			GAAGCAACAA		
Consensus	GTGCTAAGTC	AAAAAAAATC	GAAGCAACAA	GGTGTTCCGC	AGGTATCTCT
>774134IH	GGTAGAAATA				
Consensus	GGTAGAAATA	GAGTTTATAA	GTGTCAAGGA	GGCAGCTGTC	TTAGTAGAGC

# Figure 1-B

>774134IH	ATGCAGAATT	GATTCTACAA	CAAAACTGTA	TGGAAAAGAT	TGTCAATTCT
Consensus	ATGCAGAATT	GATTCTACAA	CAAAACTGTA	TGGAAAAGAT	TGTCAATTCT
>774134IH	TTCCTGATAA	AGTACAAACA	GAAAAAGCAT	CCATAATGTT	TATGCAAAGT
Consensus	TTCCTGATAA	AGTACAAACA	GAAAAAGCAT	CCATAATGTT	TATGCAAAGT
>774134IH	ATTGATTCTG	•	TTGTAACGAA		
>775437			NTGTAACGAA		
Consensus	ATTGATTCTG	TTGTTGAATT	TTGTAACGAA	AAAACCCATA	ATCAAGAAGC
>774134IH	TCCAAGCCTA	CAAAACATAA	AGTGCAATTT	TAGAAGTACA	TGGGAGGTGA
>775437	TCCAAGCCTA	CAAAACATAA	AGTGCAATTT	TAGAAGTACA	TGGGAGGTGA
Consensus	TCCAAGCCTA	CAAAACATAA	AGTGCAATTT	TAGAAGTACA	TGGGAGGTGA
>774134IH	TTAGCAATTC	TGAGGATTTT	AAAAACACCA	TACCCATGGT	GACACCACCT
>775437	TTAGCAATTC	TGAGGATTTT	AAAAACACCA	TACCCATGGT	GACACCACCT
Consensus	TTAGCAATTC	TGAGGATTTT	AAAAACACCA	TACCCATGGT	GACACCACCT
>774134IH	CCTCCACCTG	TCTTCTCATT	GCTGAAGATC	AGTCAAAGAA	TTGTGTGCTT
>775437			GCTGAAGATC		
Consensus	CCTCCACCTG	TCTTCTCATT	GCTGAAGATC	AGTCAAAGAA	TTGTGTGCTT
>774134IH	AGTTCTTGAT	AAGTCTGGAA	GCATGGGGGG	TAAGGACCGC	CTAAATCGAA
>775437	AGTTCTTGAT	AAGTCTGGAA	GCATGGGGG	TAAGGACCGC	CTAAATCGA
>1281329				TAAGGACCGC	
Consensus	AGTTCTTGAT	AAGTCTGGAA	GCATGGGGG	TAAGGACCGC	CTAAATCGAA
>774134IH	TGAATCAAGC	AGCAAAACAT	TTCCTGCTGC	AGACTGTTGA	AAATGGATCC
>1281329			TTCCTGCTGC		
Consensus			TTCCTGCTGC		
>774134IH	TGGGTGGGGA	TGGTTCACTT	TGATAGTACT	GCCACTATTG	TAAATAAGCT
>1281329			TGATAGTACT		
Consensus			TGATAGTACT		
>774134IH	ААТССАААТА	AAAAGCAGTG	ATGAAAGAAA	CACACTCATG	GCAGGATTAC
>1281329	AATCCAAATA	AAAAGCAGTG	ATGAAAGAAA	CACACTCATG	GCAGGATTAC
Consensus			ATGAAAGAAA		
>774134IH	СТАСАТАТОС	ТСТСССАССА	ACTTCCATCT	GCTCTGGAAT	TAAATATGCA
>1281329			ACTTCCATCT		
>1628677		131000100A		GCTCTGGAAT	
Consensus	CTACATATCC	TCTGGGAGGA	ACTTCCATCT		
>774134IH	TTTCAGGTGA	TTGGAGAGCT	ACATTCCCAA	CTCGATGGAT	CCGAAGTACT
>1281329	TTTCAGGTGA		-		
>1628677	TTTCAGGTGA	TTGGAGAGCT	ACATTCCCAA	CTCGATGGAT	CCGAAGTACT
Consensus	TTTCAGGTGA	TTGGAGAGCT	ACATTCCCAA	CTCGATGGAT	CCGAAGTACT

# Figure 1-C

>774134IH	GCTGCTGACT	GATGGGGAGG	ATAACACTGC	AAGTTCTTGT	ATTGATGAAG
>1628677	GCTGCTGACT	GATGGGGAGG	ATAACACTGC	AAGTTCTTGT	ATTGATGAAG
Consensus	GCTGCTGACT	GATGGGGAGG	ATAACACTGC	AAGTTCTTGT	ATTGATGAAG
>774134IH	TGAAACAAAG	TGGGGCCATT	GTTCATTTTA	TTGCTTTGGG	AAGAGCTGCT
>1628677	TGAAACAAAG	TGGGGCCATT	GTTCATTTTA	TTGCTTTGGG	AAGAGCTGCT
Consensus	TGAAACAAAG	TGGGGCCATT	GTTCATTTTA	TTGCTTTGGG	AAGAGCTGCT
>774134IH	CAMCAACCAC	maamacacam	GAGCAAGATA	1010010011	ana maaaaa
>//41341H >1628677			GAGCAAGATA		GICATITIA
Consensus			GAGCAAGATA		്നാറ മ സസസസ <b>സ</b> മ
Consensus	GAIGAAGCAG	TAATAGAGAT	GAGCAAGAIA	ACAGGAGGAA	GICAIIIIA
>774134IH	TGTTTCAGAT	GAAGCTCAGA	ACAATGGCCT	CATTGATGCT	TTTGGGGCTC
Consensus	TGTTTCAGAT	GAAGCTCAGA	ACAATGGCCT	CATTGATGCT	TTTGGGGCTC
. 77412471	mma	3 3 3 m3 cmc 3 m	OTTOTO O D O D	A CHICACOTTO A	COMOCA A A COM
>774134IH			CTCTCCCAGA		
Consensus	TTACATCAGG	AAATACTGAT	CTCTCCCAGA	AGTCCCTTCA	GCTCGAAAGT
>774134IH	AAGGGATTAA	CACTGAATAG	TAATGCCTGG	ATGAACGACA	CTGTCATAAT
>1286372					AAT
Consensus	AAGGGATTAA	CACTGAATAG	TAATGCCTGG	ATGAACGACA	CTGTCATAAT
mm 44 2 4 mm	ma1m1am1a1	00000011100	1010000000	mamax max ax	maga t a tagma
>774134IH			ACACGTTCTT		
>1286372			NCACGTTCTT		
Consensus	TGATAGTACA	GIGGGAAAGG	ACACGTTCTT	TCTCATCACA	TGGAACAGTC
>774134IH	TGCCTCCCAG	TATTTCTCTC	TGGGATCCCA	GTGGAACAAT	AATGGAAAAT
>1286372	TGCCTCCCAG	TATTTCTCTC	TGGGATCCCA	GTGGAACAAT	AATGGAAAAT
Consensus	TGCCTCCCAG	TATTTCTCTC	TGGGATCCCA	GTGGAACAAT	AATGGAAAAT
P7 44 2 4 TV	mmas as amaa	3 maa 3 3 amma	CARARMOCOC	<i>ma mome a cma</i>	mmccacca ac
>774134IH			CAAAATGGCC		
>1286372			CAAAATGGCC		
Consensus	TTCACAGTGG	A'IGCAAC'I'I'C	CAAAATGGCC	TATCTCAGTA	TTCCAGGAAC
>774134IH	TGCAAAGGTG	GGCACTTGGG	CATACAATCT	TCAAGCCAAA	GCGAACCCAG
>1286372	TGCAAAGGTG	GGCACTTGGG	CATACAATCT	TCAAGCCAAA	GCGAACCC
Consensus	TGCAAAGGTG	GGCACTTGGG	CATACAATCT	TCAAGCCAAA	GCGAACCCAG
>774134IH	አ አ አ ሮ አ ጥጥ አ አ ሮ	<i>ሞል</i> ጥጥ እር አር ጥ እ	ACTTCTCGAG	ርልርርል እ አጥጥር	ምምርም <del>ር</del> ምርርርም
	MANCALIANC	INIINCHOIN	ACTICIONG		TTCTGTGCCT
>774419					TTCTGTGCCT
>774419IH	* * * ~ * ~ * ~ *	ma.mma.o.a.coca	* 0000000000 00000 0000000000000000000		
Consensus	AAACA'I"I'AAC	TATTACAGTA	ACTTCTCGAG	CAGCAAATTC	TICIGIGCCI
>774134IH			AATGAATAAG		
>774419			ĀATGAATAAG		
>774419IH			AATGAATAAG		
Consensus			AATGAATAAG		

# Figure 1-D

>774134IH	CCCAATGATT	GTTTACGCAG	AAATTCTACA	AGGATATGTA	CCTGTTCTTG
>774419	CCCAATGATT	GTTTACGCAG	AAATTCTACA	AGGATATGTA	CCTGTTCTTG
>774419IH	CCCAATGATT	GTTTACGCAG	AAATTCTACA	AGGATATGTA	CCTGTTCTTG
Consensus	CCCAATGATT	GTTTACGCAG	AAATTCTACA	AGGATATGTA	CCTGTTCTTG
		-			
>774134IH	GAGCCAATGT	GACTGCTTTC	ATTGAATCAC	AGAATGGACA	TACAGAAGTT
>774419	GAGCCAATGT	GACTGCTTTC	ATTGAATCAC	AGAATGGACA	TACAGAAGTT
>774419IH	GAGCCAATGT	GACTGCTTTC	ATTGAATCAC	AGAATGGACA	TACAGAAGTT
Consensus	GAGCCAATGT	GACTGCTTTC	ATTGAATCAC	AGAATGGACA	TACAGAAGTT
>774134IH	TTGGAACTTT	TGGATAATGG	TGCAGGCGCT	GATTCTTTCA	AGAATGATGG
>774419	TTGGAACTTT	TGGATAATGG	TGCAGGCGCT	GATTCTTTCA	AGAATGATGG
>774419IH	TTGGAACTTT	TGGATAATGG	TGCAGGCGCT	GATTCTTTCA	AGAATGATGG
>3233118		G	TGCAGGCGCT	GATTCTTTCA	AGAATGATGG
Consensus	TTGGAACTTT	TGGATAATGG	TGCAGGCGCT	GATTCTTTCA	AGAATGATGG
>774134IH	AGTCTACTCC	AGGTATTTTA	CAGCATATAC	AGAAAATGGC	AGATATAGCT
>774419		AGGTATTTTA			
>774419IH			CAGCATATAC		
>3233118	AGTCTACTCC	AGGTATTTTA	CAGCATATAC		
Consensus	AGTCTACTCC	AGGTATTTTA	CAGCATATAC	AGAAAATGGC	AGATATAGCT
>774134IH			GGAGCAAACA		
>774419IH			GGAGCAAACA		
>3233118			GGAGCAAACA		
Consensus	TAAAAGTTCG	GGCTCATGGA	GGAGCAAACA	CTGCCAGGCT	AAAATTACGG
>774134IH			GTACATACCA		
>774419IH			GTACATACCA		TGAACGGGGA
>3233118			GTACATACCA		TGAACGGGGA
Consensus	CCTCCACTGA	ATAGAGCCGC	GTACATACCA	GGCTGGGTAG	TGAACGGGGA
				mas mas aas m	3000303003
>774134IH			GACCTGAAAT		
>774419IH		AACCCGCCAA		TGATGAGGAT	
>3233118	AA'I'I'GAAGCA	AACCCGCCAA		TGATGAGGAT	
>2733923			GACCTGAAAT		ACTCAGACCA
Consensus	AATTGAAGCA	AACCCGCCAA	GACCTGAAAT	TGATGAGGAT	ACTCAGACCA
77412471	000000000000	mmma	101001000	CACOMOCA MM	መረመረረመአምረ እ
>774134IH			ACAGCATCCG		
>774419IH			ACAGCATCCG	GAGGIGCATT	IGIGGIAICA
>3233118	CCTTGGAGGA	4		CA COMOCA MM	manaama maa
>2733923					TGTGGTATCA
Consensus	CCTTGGAGGA	TTTCAGCCGA	ACAGCATCCG	GAGGIGCATI	TGTGGTATCA
~ 77 / 1 2 / TT	C N N C C C C C C C C C C C C C C C C C	COCHHOCOCHT	GCCTGACCAA	<i>Ͳ</i> Ϡ <i>ϹϹ</i> ϹϠ <i>Ϲ</i> ϹϠϠ	CTCAAATCAC
>774134IH >774419IH					
			GCCTGACCAA GCCTGACCAA		
>2733923	CAAGTCCCAA	GCCTTCCCTT			GTCAAATNAC
>906,605	C	CCCBBCCCBB	GCCTGACCAA		
Consensus	CAAGTCCCAA	GCCTTCCCTT	GCCTGACCAA	IACCCACCAA	GICHANICAC

# Figure 1-E

>774134IH	AGACCTTGAT	GCCACAGTTC	ATGAGGATAA	GATTATTCTT	ACATGGACAG
>774419IH				GATTATTCTT	
>2733923	AGACCTTGAT	GCCACAGTTC	ATGAGGATAA	GATTATTCTT	ACATGGACAG
>906605	AGACCTTGAT	GCCACAGTTN	ATGAGGATAA	GATTATTCTT	ACATGGACAG
Consensus				GATTATTCTT	
>774134IH	CACCAGGAGA	TAATTTTGAT	GTTGGAAAAG	TTCAACGTTA	TATCATAAGA
>774419IH	CACCAGGAGA	TAATTTTGAT	GTTGGAAAAG	TTCAACGTTA	TATCATAAGA
>2733923	CACCAGGAGA	TAATTTTGAT	GTTGGAAAAG	TTCAACGTTA	TATCA
>906605	CACCAGGAGA	TAATTTTGAT	GTTGGAAAAG	TTCAACGTTA	TATCATAAGA
Consensus	CACCAGGAGA	TAATTTTGAT	GTTGGAAAAG	TTCAACGTTA	TATCATAAGA
>774134IH	ATAAGTGCAA	GTATTCTTGA	TCTAAGAGAC	AGTTTTGATG	ATGCTCTTCA
>774419IH	ATAAGTGCAA	GTATTCTTGA	TCTAAGAGAC	AGTTTTGATG	ATGCTCTTCA
>906605	ATAAGTGCAA	GTATTCTTGA	TCTAAGAGAC	AGTTTTGATG	ATGCTCTTCA
>2771475	AA	GTATTCTTGA	TCTAAGAGAC	AGTTTTGATG	ATGCTCTTCA
Consensus	ATAAGTGCAA	GTATTCTTGA	TCTAAGAGAC	AGTTTTGATG	ATGCTCTTCA
>774134IH				GGCCAACTCC	
>774419IH				GGCCAACTCC	
>906605				GGCCAACTCC	
>2771475				GGCCAACTCC	
Consensus	AGTAAATACT	ACTGATCTGT	CACCAAAGGA	GGCCAACTCC	AAGGAAAGCT
>774134IH				AAAATGCAAC	
>774419IH				AAAATGCAAC	
>906605				AAAATGCAAC	
>2771475	TTGCATTAA	ACCAGAAAA'I'		AAAATGCAAC	
>1803247	mma a 2 mmm 2 2	1001011110		ANAATGCAAC	
Consensus	TIGCATITAA	ACCAGAAAAT	ATCTCAGAAG	AAAATGCAAC	CCACATATTI
>774134IH	<b>አጥጥ</b> ርርር እጥጥ አ	AAAGTATAGA	ТАЛААССААТ	TTGACATCAA	AAGTATCCAA
>774419IH				TTGACATCAA	
>906605				TTGGCATCAA	
>2771475				TTGACATCAA	
>1803247				TTGACATCAA	
Consensus	ATTGCCATTA	AAAGTATAGA	TAAAAGCAAT	TTGACATCAA	AAGTATCCAA
>774134IH	CATTGCACAA	GTAACTTTGT	TTATCCCTCA	AGCAAATCCT	GATGACATTG
>774419IH	CATTGCACAA	GTAACTTTGT	TTATCCCTCA	AGCAAATCCT	GATGACATTG
>2771475	CATTGCACAA	GTAACTTTGT	TTATCCCTCA	AGCAAATCCT	GATGACATTG
>1803247	CATTGCACAA	GTAACTTTGT	TTATCCCTCA	AGCAAATCCT	GATGACATTG
Consensus				AGCAAATCCT	
			_		
>774134IH				ATAAAAGTCA	
>774419IH				ATAAAAGTCA	
>1803247	ATCCTACTCC			ATAAAAGTCA	
>1737526					TAATTCTGGA
Consensus	ATCCTACTCC	TACTCCTACT	CCTACTCCTG	ATAAAAGTCA	TAATTCTGGA

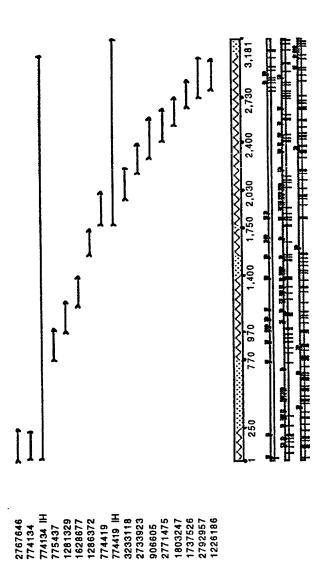
### Figure 1-F

>774134IH			ATTGTCTGTG		
>774419IH			ATTGTCTGTG		TTGTAATTGT
>1803247			ATTGTCTGTG		
>1737526	GTTAATATTT	CTACGCTGGT	ATTGTCTGTG		
>2792957					TTGTAATTGT
Consensus	GTTAATATTT	CTACGCTGGT	ATTGTCTGTG	ATTGGGTCTG	TTGTAATTGT
>774134IH	TAACTTTATT	TTAAGTACCA	CCATTTGAAC	CTTAACGAAG	АААААААТСТ
>774419IH	TAACTTTATT	TTAAGTACCA	CCATTTGAAC	CTTAACGAAG	AAAAAAATCT
>1737526	TAACTTTATT	TTAAGTACCA	CCATTTGAAC	CTTAACGAAG	АААААААТСТ
>2792957	TAACTTTATT	TTAAGTACCA	CCATTTGAAC	CTTAACGAAG	AAAAAAATCT
>1226186				-	AAAAAAATCT
Consensus	TAACTTTATT	TTAAGTACCA	CCATTTGAAC	CTTAACGAAG	AAAAAAATCT
>774134IH	TCAAGTAGAC	CTAGAAGAGA	GTTTTAAAAA	ACAAAACAAT	GTAAGTAAAG
>774419IH			GTTTTAAAAA		GTAAGTAAAG
>1737526	TCAAGTAGAC	CTAGAAGAGA	GTTTTAAAAA	ACAAAACAAT	GTAAGTAAAG
>2792957	TCAAGTAGAC	CTAGAAGAGA	GTTTTAAAAA	ACAAAACAAT	GTAAGTNAAG
>1226186			GTTTTAAAAA		GTAAGTAAAG
Consensus	TCAAGTAGAC	CTAGAAGAGA	GTTTTAAAAA	ACAAAACAAT	GTAAGTAAAG
BB 44 3 4 TV		አ አ ጠረመመን አ አ አ	TTCATCCCAT	стстсатсат	AAACTCATAA
>774134IH	CATATTICIG	AMICIIAAAA	TTCATCCCAT	GTGTGATCAT	AAACTCATAA
>774419IH >1737526	GATATTTCTG		HEATCOM	01010111	
			TTCATCCCAT	GTGTGATCAT	AAACTCATAA
>2792957 >1226186	CATATITUTE	አልተርተገለለለ	TTCATCCCAT	GTGTGATCAT	AAACTCATAA
	CATATTICIG	አለጥርጥጥል እል	TTCATCCCAT	GTGTGATCAT	AAACTCATAA
Consensus	GAIAIIICIG	ARICITATION	110/11000111		
>774134IH	AAATAATTTT	AAGATGTCGG	AAAAGGATAC	TTTGATTAAA	TAAAAACACT
>774419IH			AAAAGGATAC	TTTGATTAAA	TAAAAACACT
>2792957			AAAAGGATAC	TTTGATTAAA	TAAAAACACT
>1226186	TTTTAATAAA	AAGATGTCGG	AAAAGGATAC		TAAAAACACT
Consensus	AAATAATTTT	AAGATGTCGG	AAAAGGATAC	TTTGATTAAA	ТАААААСАСТ
>774134IH			CAAGATTAAA		
>774419IH			CAAGATTAAA		
>2792957	CATGGATATG	TAAAAACTGT	CAAGATTAAN	ATTTAATAGT	TTCATTTATT
>1226186	CATGGATATG	TAAAAACTGT	CAAGATTAAA	ATTTAATAGT	T"I'CAT"I"TAT"I
Consensus	CATĠGATATG	TAAAAACTGT	CAAGATTAAA	ATTTAATAGT	TTCATTTATT
>774134IH	TGTTATTTA	TTTGTAAGAA	ATAGTGATGA	ACAAAGATCC	TTTTTCATAC
>774419IH	TGTTATTTTA	TTTGTAAGAA	ATAGTGATGA	ACAAAGATCC	TTTTTCATAC
>2792957			ATAGTGATGA		
>1226186	TGTTATTTT	TTTGTAAG			
Consensus	TGTTATTTT	TTTGTAAGAA	. ATAGTGATGA	ACAAAGATCC	TTTTTCATAC
	ma3.m	-			
>774134IH	TGAT			<u>് സ്ഥാനന്നാന് സ്ഥാന</u> ്	AAATGATATT
>774419IH	TGATACCTGG	TIGTATAT	TITIGATGCAA	CAGIIIICIG CAGIIIICIG	AAATGATATT
Consensus	TGATACCTGC	5 TTGTATATT	TITGATGCAA	CAGITICIE	i multipliant
>774419IH	TCAAATTGCA	TCAAGAAATT	AAAATCATCT	ATCTGAGTAG	TCAAAATACA
Consensus	TCAAATTGCA	A TCAAGAAATT	AAAATCATCI	ATCTGAGTAG	TCAAAATACA

### Figure 1-G

>774419IH Consensus AGTAAAGGAG AGCAAATAAA CAACATTTGG A AGTAAAGGAG AGCAAATAAA CAACATTTGG A

Figure 2



12.2% Halling